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December 13, 2002, 15:41:52; Search time 268 Seconds (Without alignments) 1378.089 Million cell updates/sec
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164
1 gcaccgggacatcaaggcag......tgcctggatggccccagaag 164
GenCore version 5.1.3 Copyright (c) 1993 - 2002 Compugen Ltd.
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Listing first 45 summaries
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES ID Descrip	AAF30487	AAH46913	ABQ86165	ABN86357 Novel human protei	ABN86358	AAD18824	AAD34309	AA185458	
DB	22	22	24	24	24	22	. 24	22	
% Query Match Length DB	164	2157	3066	3111	3518	3141	3538	444	6
& Query Match	100.0	93.2	93.2	93.2	93.2	57.4	57.4	53.0	,
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Vulnerary; ss. Homo sapiens. Key CDS EP1085093-A2. 21-MAR-2001, 12-SEP-2000; 20 20-SEP-1999;	mmatory; antipsoriatic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Protein Kinase; enzyme; cytostatic; nootropic; neuroprotective; human, antiparkinsonian; virucide; antibacterial; antifungal; antimigraine; analyesic; hypotensive; immunosuppressive; antiallergic; antipheumatic; antiarthritic; ophthalmological; anorectic; osteopathic; thrombolytic; antiateriosclerotic; antiasthmatic; vasotropic; antidiabetic; gene therapy; ss.
                                                                                                                                                                                               The present sequence is that of a partial CDNA encoding MLK4 (see AAB20333), 104 novel c-Jun N-terminal kinase kinases constants of the invention. CDNAs encoding JNKKKS were isolated from human foreskin epidermal Aeratinocyte cDNA by PCR amplification using primars (see AAB30494-97) based on regions of homology between the kinase domains of JNKKS or JNKKKS. MLK4 was 50-78% identical to members of the MLK family of JNKKKS. The MLK4 gene is cappeased in Keratinocytes, kidney and pancreas, but not in brain, placenta, lung, liver or skeletal muscle. The transcript size is products are useful for elucidation of the components involved in the cellular response to ultraviolet radiation. They can be used in drug discovery, by screening for compounds that affect the collular JNKKK. Particularly useful are drugs that reduce UV according a JNKKK. Particularly are drugs that reduce UV
                                                                                         New polynucleotides encoding c-Jun N-terminal kinase kinases kinases in C. MIKM, PAKM, associated with skin damage for use in drug screening and development.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 GCACCGGGACATCAAGGCAGGAAATATTTGCTACTTGAGAAGATAGAACATGATGACAT 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ouery Match 100.0%; Score 164; DB 22; Length 164; Best Local Similarity 100.0%; Pred, No. 19e-45; Matches 164; Conservative 0; Mismatches 0; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      121 CAAAATGAGCACAGCACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           121 CAAAATGAGCACAGCAGCACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 164 BP; 55 A; 33 C; 44 G; 32 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cDNA encoding human protein kinase SGK067.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        drugs that enhance wound healing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAH46913 standard; cDNA; 2157 BP.
                                                                                                                                                                 Claim 3; Page 22; 51pp; English.
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Blumenberg M, Gazel AM;
                                 WPI; 2001-236883/25.
                                                       P-PSDB; AAB20333
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continuous provinces numeral process and process. The kinase corrections and polymetrides encoding the polymetrides. The kinase colleges and their modulators are useful for treating a disease or disorder such as cancer, immune-related diseases, cardiovascular disease, brain or neuronal associated disease and metabolic disorders, including cancers of tissues, cancers of hematopoletic origin, diseases of the cancers of tissues, cancers of hematopoletic origin, diseases of the cancers of tissues, viral infections, infections caused by prions, lateral sclerosis, viral infections, infections caused by prions, bacteria and fungi, ocular diseases, multiple sclerosis, amyotrophic lateral sclerosis, viral infections, infections caused by prions, mod disorders, attention disorders, cognition disorders, hypotension, hypotension, psychotic disorders, cognition disorders, hypotension, hypotension, psychotic disorders, neurological disorders, hypotension, metabolic disorders, autoimmunity, atherosclerosis, psoriasis, correction citianmatory bowel disease, rhemmatory pelvic disease, chronic infinammatory pelvic disease, chronic infinammatory pelvic disease, chronic infinammatory belvic diseases such as reperfusion injury, cornoary thrombosis, clotting disorders and atherosclerosis, coular diseases such as glaucoma, retinopathy and macular degeneration, psychiatric and neurological disorders such as anxiety, scholaring, psychiatric and neurological disorders such as anxiety, scholaring, and any prince of the collar diseases such as allocand, retinopathy as anxiety or metabolic disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                dementia, manic depression, etc. The polynucleotides are useful in gene therapy techniques to treat the above mentioned disorders. Sequences AAH46891-46922 represent human protein kinases encoding cDNA molecules.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61 CTGCAATAAAACTTTGAAGATTACAGATTTTGGGTTGGCGAGGGAATGGCACAGGACCAC 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Khinase polypeptides useful for treating cancers, Alzheimer's disease, viral infections, diabetes, obesity, organ transplant rejection and rheumatoid arthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention provides human protein kinases and protein kinase-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Plowman G, Whyte D, Manning G, Sudarsanam S, Martinez R;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Page 209-210; 218pp; English.
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                                                                                                                                                                                                                                   2000US-0190162.
2000US-0193404.
25-JAN-2001; 2001WO-US02337.
                                                                                           2000US-0178078.
                                                                                                                                       2000US-0179364.
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                                                                                                                                                                                                                                                                                                                                                                                                                                (SUGE-) SUGEN INC.
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                                                                                                                                 31-JAN-2000;
17-FEB-2000;
17-MAR-2000;
29-MAR-2000;
13-NOV-2000;
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                                                                                           Human; cytostatic; vulnerary; antiarteriosclerotic; antiparkinsonian;
M nottropic: neuroprotective; immunosuppressive; haemostatic;
M antiinflammatory; cardiant; antiulcer; virucide; antithyroid;
cerebroprotective, anorectic; metabolic; vaccine; cancer; infection;
wound healing disorders; atherosclerosis; Parkinson's disease;
M Alcheimer's disease; autoimmune disorder; haematopoietic disorder;
Inflammation; neoplastic disoase; nervous system disorder;
Inflammation; neoplastic disease; nervous system disorder;
Archeimer disorders; pancreatitis; respiratory disorder;
M perproliferation; systemic autoimmune disease; hyper-immunity;
developmental abnormality; gastrointestinal ulceration; neuropathy;
M haematological disease; sperm dysfunction;
thyroid disorder; hypothyroidism; brain damage; colitis;
cone photo- transduction deficiency; neurological disease; stroke;
M anglogenesis; ovulation disorder; spinal cord; thyroid gland; heart;
M rachea; thymus; lymph node; muscular system; obesity; anorexia;
M growth abnormality; precocious puberty; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Secreted proteins and polynucleotides useful as vaccines for preventing or treating various diseases e.g. cancer, wounds, atherosclerosis, parkinson's disease, Alfalamer's disease, infection, autolumune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cogswell JP, Kabnick KF, L
Smith RF, Strum JC, Xie Q;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BEECHAM PLC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-JAN-2001; 20010S-260482P.
30-JAN-2001; 20010S-26492P.
66-FEB-2001; 20010S-26797P.
19-MAR-2001; 20010S-27698BP.
04-APR-2001; 2001US-281535P.
                                                     Novel human gene. SEQ ID 36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-DEC-2001; 2001WO-US49232.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2000US-256710P.
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10-SEP-2002 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GLAX ) GLAXO GROUP LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2002-508784/54.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Agarwal P,
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trachea, thymus, lymph node and muscular system, obesity, anorexia, growth abnormalities, and alleviation of precocious puberty. The sequences given in records ABQ86130-ABQ86184 represent novel human cDNA's of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                          61 CTGCAATAAAACTTTGAAGATTACAGATTTTGGGTTGGCGAGGGAATGGCACAGAGACCAC 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 GCACCGGGACATCAAGGCAGGAAATATTTTGCTACTTGAGAAGATAGAACATGACAT 60
                                                                                                                                                                                                                                      93.2%; Score 152.8; DB 24; Length 3066; 95.7%; Pred. No. 4e-41; 1ve 0; Mismatches 7; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /*tag= a
/transl_except= (pos: 2773..2775, aa: Xaa)
/product= "NHP kinase"
                                                                                                                                                                           Sequence 3066 BP; 738 A; 881 C; 852 G; 595 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         121 CAAAATGAGCACAGCAGCACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              855 CAAAATGAGCACAGCAGCACCTATGCCTGGATGGCCCCCGGAG 898
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel human protein; NHP; kinase; human; gene; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel human protein (NHP) kinase coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 36-37; 41pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and pharmacogenomic applications -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABN86357 standard; DNA; 3111 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (LEXI-) LEXICON GENETICS INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-OCT-2002 (first entry)
                                                                                                                                                                                                                                                                                                         Matches 157; Conservative
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P-PSDB; ABB80923.
                                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                               Best Local
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Human kinase; PKIN; gene therapy; adenocarcinoma; immune disorder; gout; cancer; allergy; sarcoma; leukaemia; acquired immune deficiency syndrome; AIDS; Addison's disease; mircobial infection; inflammation; osteoporosis; atherosclerosis; cardiovascular disease; myocardial infarction; anaemia; myasthenia gravis; cirrhosis; cattaract; growth and development disorder; selzure disorder; pulmonary embolism; Gaucher's disease; lipid disorder; lipid storage disease; pick's disease; Tay-Sachs disease; renal disease; obesity; restorative therapy; immunomodulatory; vaccine; cardiovascular; antimicrobial; cytostatic; antiliflammatory; asthma; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gandhi AR, Tribouley CM, Kearney L, Griffin JA, Nguyen DB; O, Lu DAM, Lal P, Burford N, Khan FA, Walia NK, Yao MG; On C, Burrill JD, Marcus GA, Zingler KA, Recipon SA, Lu Y; JL, Thornton M, Tang YT, Hafalia A, Elliott VS, Baughn MR; T, Ramkumar J, Borowsky ML, Au-young J, Hillman JL;
                         61 CTGCAATAAAACTTTGAAGATTACAGATTTTGGGTTGGCGAGGGAATGGCACAGGACCAC 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human kinases and nucleic acids, useful for preventing diagnosing and treating cancers, inflammation and immune disorders -
  1 GCACCGGGACATCAAGGCAGGAAATATTTTGCTACTTGAGAAGATAGAACATGATGACAT 60
                                                                                                                                                                                    1163 CAAAATGAGCACACGAGCACCTATGCCTGGATGGCCCCCGAAG 1206
                                                                                                                                                                121 CAAAATGAGCACAGGCACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /product= "Human PKIN-9 protein"
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                                                                                                                                                                                                                                                                                                           AAD18824 standard; cDNA; 3141 BP.
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28-APR-2000; 2000US-200226P.
05-MAY-2000; 2000US-203339P.
11-MAY-2000; 2000US-203505P.
18-MAY-2000; 2000US-205564P.
26-MAY-2000; 2000US-2075739P.
01-JUN-2000; 2000US-208795P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Human kinase (PKIN)-9 cDNA.
                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2001-611740/70.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200181555-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Walsh RT, R
Gururajan R;
                                                                                                                                                                                                                                                                                                                                                                                            18-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Policky JL,
Walsh RT, R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-NOV-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bandman O,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patterson
                                                                                                                                                                                                                                                                                                                                                    AAD18824;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Yue H,
                                                                                                                                                                                                                                                                                      AAD18824
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a novel human protein (NHP), kinase that shares structural similarity with animal kinases. The kinase polynucleotides are useful in therapeutic, diagnostic and pharmacogeomonic applications and for identifying compounds that modulate, i.e. act as agonists or antagonists of the gene expression or gene product activity. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel human kinase polynucleotide encoding a protein that shares structural similarity with animal kinases for therapeutic, diagnostic
                                                                                                                                               CTGCAATAAAACTTTGAAGATTACAGATTTTGGGTTGGCGAGGGAATGGCACAGGACCAC 120
                                                                                                                                                                                                                            0; Gaps
                                                                                                                      1 GCACCGGGACATCAAGGCAGGAAATATTTTGCTACTTGAGAAGATAGAACATGATGACAT
                                  93.2%; Score 152.8; DB 24; Length 3111; ilarity 95.7%; Pred. No. 4.1e-41; Conservative 0; Mismatches 7; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          93.2%; Score 152.8; DB 24; Length 3518; 95.7%; Pred. No. 4.3e-41; trive 0; Mismatches 7; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 3518 BP; 811 A; 1043 C; 976 G; 680 T; 8 other;
Sequence 3111 BP; 740 A; 905 C; 864 G; 594 T; 8 other;
                                                                                                                                                                                                                                                                                                             900 CAAAATGAGCACAGGGACCTATGCCTGGATGGCCCCCGAAG 943
                                                                                                                                                                                                                                                                                    121 CAAAATGAGCACAGCAGCACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel human protein; NHP; kinase; human; gene; ds.
                                                   4.1e-41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequence represents the NHP kinase encoding DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel human protein (NHP) kinase encoding DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 39-41; 41pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
264..3374
/*tag= a
/product= "kinase"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and pharmacogenomic applications -
                                                                                                                                                                                                                                                                                                                                                                                                                               ABN86358 standard; DNA; 3518 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (LEXI-) LEXICON GENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-DEC-2000; 2000US-254744P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "NHP"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-DEC-2001; 2001WO-US47606.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hu Y, Kieke JA, Donoho G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-OCT-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 95.7
Matches 157; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-566739/60.
                                                     Local Similarity
es 157; Conservi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200255685-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-JUL-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABN86358:
                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                            datches
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The present invention relates to human kinases (PKIN) and the nucleic acids encoding them. PKIN is used as vaccine and in gene therapy. PKIN is

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Gaps

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used in the prevention, diagnosis and treatment of diseases cancers, adenocarcinoma, leukaemia, sarcoma, immune disorder, Addison's disease, acquired immune deficiency syndrome (AIDS), anaemia, asthma, allergies, gout, microbial infections, cardiovascular disease and/or inflammation, myasthenia gravis, atheroscolerosis, cirrhosis, osteoporosis, myocardial, myasthenia gravis, atheroscolerosis, cirrhosis, osteoporosis, myocardial, infarction, cataract, growth and development disorder, seizure disorder, cinfarction, cataract, growth and development disorder, lipid storage disease, pick's disease, pray-Sachs disease, renal disease and obesity.

Expression by rectifying mutations or deletions in a patient's genome that affect the activity of PXIN by expressing inactive proteins or to supplement the patients own production of PXIN nucleic acids may be used to produce the PXIN polypeptide, by inserting the nucleic acids may be used to recoduce the PXIN polypeptide, by inserting the nucleic acids into a host cell and culturing the cell to express the protein. PXIN nucleic acid sequences in samples and therefore which patients may be in need of restorative therapy. The present sequence is human PXIN-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        acquired immune deficiency syndrome; thymic hypoplasia; Crohn's disease; asthma; neurological disorder; epilepsy; Charcot-Marie-Tooth disease; AIDS; selzures; cell proliferative disorder; cancer; adenocarcinoma; leukaemia; lymphoma; melanoma; myethoma; sarcoma; developmental disorder; Down's syndrome; gene therapy; protein therapy; cytostatic; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; kinase; enzyme; PKIN-12 protein; immune system disorder; anaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          62 TGCAATAAAACTTTGAAGATTACAGATTTTGGGTTGGCGAGGGAATGGCACAGGACCACC 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 CACCGGGACATCAAGGCAGGAAATATTTTGCTACTTGAGAAGATAGAACATGATGACATC 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 57.4%; Score 94.2; DB 22; Length 3141; Best Local Similarity 73.6%; Pred. No. 2.3e-21; Matches 120; Conservative 0; Mismatches 43; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /product= "Human mature PKIN-12 protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 3141 BP; 685 A; 941 C; 942 G; 573 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   122 AAAATGAGCACAGCAGCACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /product= "Human PKIN-12 protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAD34309 standard; cDNA; 3538 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-AUG-2001; 2001WO-US27219.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-JUL-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ø
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52..3291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human PKIN-12 cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200218557-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-MAR-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mat_peptide
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AAD34309
ID AAD34309
ID AAD34309
DT 16-2
XX
XX
XX
KW HUME
KW AIDS
KW 
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The present invention relates to human kinases (PKIN) and polynucleotides encoding such proteins. PKIN sequences of the invention are useful for diagnosing, treating or preventing disorders associated with aberrant expression of PKIN, particularly immune system disorders (e.g. acquired immune deficiency syndrome (AIDS), thymic hypoplasta, Crohn's disease, an emmia, asthma), neurological disorders (e.g. epilepsy, Charcot-Marier Tooth disease or seizures), cell proliferative disorders (e.g. cancers such as adenocarcinoma, leukaemia, lymphoma, melanoma, melanoma, sarcoma, and developmental disorders (e.g. Down's syndrome). They are also used in gene therapy and protein therapy. The present sequence is a CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; cytokine; cell proliferation; cell differentiation; gene therapy; vaccine; peptide therapy; stem cell growth factor; hematopolesis; tissue growth factor; immunomodulatory; cancer; leukaemia; nervous system disorders; arthritis; inflammation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                         New human kinases, useful for diagnosing, treating or preventing immune system disorders (e.g. Crohn's disease), neurological disorders (e.g. epilepsy), or cell proliferative disorders (e.g. cancers such as leukemia or lymphoma)
                                                                                                                                                                                                                                             Tribouley CM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    62 TGCAATAAAACTTTGAAGATTACAGATTTTGGGTTGGCGAGGGAATGGCACAGGACCACC 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 CACCGGGACATCAAGGCAGGAAATATTTTGCTACTTGAGAAGATAGAACATGATGACATC 61
                                                                                                                                                                                                                         Gandhi AR;
                                                                                                                                                                                                                                                                                   PG;
                                                                                                                                                                                                                       Nguyen DB, Walia NK, Hafalia AJA, Yao MG, Gandhi
Ding L, Patterson C, Yue H, Baughn MR, Triboule
Elliott VS, Lu Y, Ison CH, Au-Young J, Tang YT;
Burrill JD, Marcus GA, Zingler KA, Lu DAM, Lal PG
Warren BA, Kearney L, Policky JL, Thangavelu K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 57.4%; Score 94.2; DB 24; Length 3538; Best Local Similarity 73.6%; Pred. No. 2.4e-21; Matches 120; Conservative 0; Mismatches 43; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 3538 BP; 763 A; 1055 C; 1062 G; 658 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  122 AAAATGAGCACAGCAGCACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 91; Page 207-208; 218pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human polynucleotide SEQ ID NO 5518.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAI85458 standard; cDNA; 444 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           encoding human PKIN-12 protein.
              31 AUG 2000; 2000US 229973P.

08-SEP-2000; 2000US 231357P.

22-SEP-2000; 2000US 23490ZP.

25-SEP-2000; 2000US 23490ZP.

06-OCT-2000; 2000US -236499P.
                                                                                                                                             13-OCT-2000; 2000US-240542P.
                                                                                                                                                                                   (INCY-) INCYTE GENOMICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-NOV-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-329769/36.
P-PSDB; AAE21717.
                                                                                                                                                                                                                                                                  Thornton M,
Azimzai Y, B
Ramkumar J,
                                                                                                                                                                                                                                                 Gururajan R,
                                                                                                                                                                                                                              Bandman O,
                                                                                                                                                                                                                                                                                                                                   Burford N;
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AA1854584
ID AA18541
XX AA2 AA18541
XX XX AA18541
XX XX AA18541
XX XX WW HUMBU 1
XX W WACCING
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WO200228999-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to human polynucleotides (AAI79941-AAI93841) and the encoded proteins (AA000010-AA019910) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The polynelectides and polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypetides have various cytokine-like activities, e.g. stem cell growth factor activity, haematopolesis regulating activity tissue growth factor activity, immunomodulatory activity and activity and may be useful in the diagnosis and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO and fifth wipo.int/pub/published_pot_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           treatment of cancer, leukaemia, nervous system disorders, arthritis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    74 TIGAAGATTACAGATTITGGGTTGGCGAGGGAATGGCACAGGACCACCAAAATGAGCACA 133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           443 AAGTCCAGCAACATATTGATCCTCCAGAAGGTGGAGAATGGAGACCTGAGCAACAAGATT 384
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14 AAGGCAGGAAATATTTGCTACTTGAGAAGATAGAACATGATGACATCTGCAATAAAACT 73
                                                                                                                                                                                                                                                                                                                                                Isolated nucleic acids and polypeptides, useful for preventing diagnosing and treating e.g. leukaemia, inflammation and immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human cDNA differentially expressed in granulocytic cells #445.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human, ss, granulocytic cell; DNA chip; bacterial infection; viral infection; protozoal infection; fungal infection; sterile inflammatory disease; psoriasis; rheumatoid arthritis; glomerulonephritis; asthma; thrombosis;
                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO 5518; 1399pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     53.0%; Score 87; DB 22; Length 444; 73.5%; Pred. No. 2.7e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             40; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 444 BP; 110 A; 119 C; 111 G; 101 T; 3 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   323 GCAGGGACGTATGCTTGGATGGCACCCGAAG 293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            134 GCAGGCACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABK83874 standard; cDNA; 3558 BP.
                                                                                                                                                                                                                                                          Tang YT, Liu C, Drmanac RT;
                                                                                                                         26-FEB-2001; 2001WO-US04927.
                                                                                                                                                             28-FEB-2000; 2000US-0515126
18-MAY-2000; 2000US-0577409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-AUG-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches 111; Conservative
                                                                                                                                                                                                                                                                                         WPI; 2001-514838/56.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                      (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                              P-PSDB; AAO05527
                                                  WO200164835-A2.
                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                inflammation
                                                                                        07-SEP-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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The invention relates to detecting (MI) granulocyte (GC) activation (CCA), by defecting the level of expression of gene(s) (GS) identified by DNA chip analysis as given in the specification, and comparing the expression level in an unactivated comparing the expression level in an unactivated (GC, where differential expression of GS is indicative of GA. Also included are modulating (M2) GA. by contacting GC with an agent capable of modulating GA. by contacting GC with an agent capable of modulating GA. by contacting GS (2) screening (M3) for an agent capable of modulating GA or an inflammation (especially chronic) in a tissue, an allergic response in a subject, exposure of a subject to a pathogen or sterile inflammatory disease using the crossiblect to a pathogen or sterile inflammatory disease using the crossiblect to a pathogen or sterile inflammatory disease. by detecting the crossiblect to a pathogen or sterile inflammatory disease, by detecting the crossiblect contacting contacting the crossiblect contacting contacting a tissue having contacting the crossiblect contacting contacting crossiblect contacting contacting contacting crossiblect contacting 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detecting granulocyte activation by detecting differential expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 CACCGGGACATCAAGGCAGGAAATATTTTGCTACTTGAGAAGATAGAACATGATGACATC 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of genes associated with granulocyte activation, which serves as diagnostic markers that is useful for monitoring disease states and
cardiac reperfusion injury; renal reperfusion injury; ARDS; adult respiratory distress syndrome; inflammatory bowel disease; Crohn's disease; ulcerative colitis; periodontal disease; granulocyte activation; chronic inflammation; allergy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 3558;
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69.3%; Pred. No. 1.4e-17;
Live 0; Mismatches 50; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 3558 BP; 635 Å; 1211 C; 1161 G; 551 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Beazer-Barclay Y, Weissman SM, Yamaga S, Vockley J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; SEQ ID No 445; 114pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-OCT-2001; 2001WO-US30821.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-OCT-2000; 2000US-237189P.
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Matches 113; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GENE-) GENE LOGIC INC.
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Human; cancer; colon; breast; ovary; oesophagus; kidney; thyroid;
stomach; lung; prostate; pancreas; carcinoma; antiumour; cancerous;
cytostatic; gene therapy; antineoplastic; wllm's tumour; adenocarcinoma;
                                                                                                                                                                               62 TGCAATAAAACTTTGAAGATTACAGATTTTGGGTTGGCGAGGGAATGGCACAGGACCACC 121
                                                                                                         Gaps
                                                                                                                                                    2 CACCGGGACATCAAGGCAGGAAATATTTTGCTACTTGAGAAGATAGAACATGATGACATC 61
                                                                                                         ö
                                                              Length 3558;
                                                        Query Watch 50.6%; Score 83; DB 24; Length 35; Best Local Similarity 69.3%; Pred, No. 1.4e-17; Matches 113; Conservative 0; Mismatches 50; Indels
             Sequence 3558 BP; 635 A; 1211 C; 1161 G; 551 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                               Pancreas cancer related gene sequence SEQ ID NO:8355.
                                                                                                                                                                                                                                                                                                                                    122 AAAATGAGCACAGCAGCACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABL70018 standard; DNA; 3454 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           200005-234009P.
200005-234034P.
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29-SEP-2000;
02-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABL70018;
                                                                                                                                                                                                                                                                                      qq
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a method for screening biologically active agent that modulates cancer associated protein kinase function. The invention also relates to a method for diagnosing cancer comprising determining the upregulation of expression of a nucleic acid sequence encoding a protein upregulation of expression of a nucleic acid sequence encoding a protein to the method is useful for screening biological agents that modulate cancer associated correction kinase function. Downregulating the activity of protein kinase is useful for inhibiting the growth of a cancer cell, e.g. liver or colon cancer. A nucleic acid encoding protein kinase is useful to screen biopsy cancer. A nucleic acid encoding protein kinase is useful to screen biopsy cancer. A nucleic and inflammatory samples such as arthritic synovium, for amplified DNA in the cell or increased expression of corresponding mRNA or protein and is also useful to detect differences in expression levels cancer as only as molecular weight, amino acid and nucleotide sequences between the cancer cells. The present sequence is human mitogen activated protein kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; cytostatic; antisense gene therapy; screening; protein kinase; cancer; liver; colon; tumour; inflammation; arthritic synovium; MAP3K11; mitogen activated protein kinase; enzyme; gene; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Diagnosing cancer, comprises determining the upregulation of expression of a nucleic acid sequence encoding a protein kinase or upregulation of expression of the protein kinase, in the cancer
1196 CACCGTGATCTCAAGTCCCAACAACATTTTGCTGCTGCAGCCCATTGAGAGTGACGACATG 1255
                                                                62 TGCAATAAAACTTTGAAGATTACAGATTTTGGGTTGGCGAGGGAATGGCACAGGACCACC 121
                                                                                                                                                       Human mitogen activated protein kinase, MAP3K11 gene.
                                                                                                                                 122 AAAAIGAGCACAGCAGCACCTAIGCCTGGAIGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /product= "Human MAP3K11 protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 16; Page 56-60; 87pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                         AAD36139 standard; DNA; 3558 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (KINE-) KINETEK PHARM INC.
(UYBR-) UNIV BRITISH COLUMBIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-SEP-2000; 2000US-233999P. 02-CCT-2000; 2000US-337419P. 04-CCT-2000; 2000US-237423P. 04-CCT-2000; 2000US-29555P. 10-MAY-2001; 2001US-290555P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-SEP-2001; 2001WO-IB02237.
                                                                                                                                                                                                                                                                                                                                                                               09-AUG-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Yoganathan I, Delaney AD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-394145/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              P-PSDB; AAE22763.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200224947-A2.
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                                                                                                                                                                                                                                                                                                                                       AAD36139;
                                                                                                                                                                                                                                           RESULT 10
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31-JAN-2000; 2000US-0119065.

24-FEB-2000; 2000US-0186528.

25-FEB-2000; 2000US-01865350.

26-MAR-2000; 2000US-01861351.

19-MAR-2000; 2000US-0198131.

19-MAR-2000; 2000US-0198131.

19-MAR-2000; 2000US-0198131.

19-MAR-2000; 2000US-019813.

07-JUL-2000; 2000US-0218486.

11-JUL-2000; 2000US-0218487.

11-JUL-2000; 2000US-0218487.

11-JUL-2000; 2000US-0218487.

11-JUL-2000; 2000US-0218487.

11-JUL-2000; 2000US-0218487.

11-JUL-2000; 2000US-0218487.

11-JUL-2000; 2000US-0218497.

11-JUL-2000; 2000US-0218296.

26-JUL-2000; 2000US-0218296.

26-JUL-2000; 2000US-0218296.

26-JUL-2000; 2000US-0228287.

14-MG-2000; 2000US-0228287.

14-MG-2000; 2000US-0228287.

14-MG-2000; 2000US-0228288.

14-MG-2000; 2000US-0228287.

14-MG-2000; 2000US-0228287.

14-MG-2000; 2000US-0228289.

14-MG-2000; 2000US-0228289.

14-MG-2000; 2000US-0228287.

14-MG-2000; 2000US-0228289.

16-SEP-2000; 2000US-0228144.

16-SEP-2000; 2000US-0228148.

16-SEP-2000; 2000US-0228148.

16-SEP-2000; 2000US-023148.

16-SEP-2000; 2000US-023148.

16-SEP-2000; 2000US-0231398.

16-SEP-2000; 2000US-0231398.

11-SEP-2000; 2000US-0231399.

11-SEP-2000; 2000US-0231399.
                                                                                                                                                                                                                                       17-JAN-2001; 2001WO-US01334
                                                                                                                                                                    WO200159063-A2.
                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                       16-AUG-2001.
ZZZZZZZ
                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes a method (M1) for screening for an anti-nephastic agent. The method involves exposing cells to a chemical agent to be tested for anti-nephastic activity, determining a change in expression of at least one gene (I) of a signature gene set, where (I) comprises a sequence (S) selected from 844 7 sequences (given in ABLO1664 to ABL70110), or is at least 95% identical to (S), where a change in expression is indicative of anti-neoplastic activity. (I) has oytostatic activity and can be used in gene therapy. M1 can be used for screening an anti-neoplastic agent, and can be used for producing a product which is the data collected with respect to the anti-neoplastic agent as a result of M1, and the data is sufficient to convey the chemical structure and/or properties of the agent. M1 can be used in the treatment of cancer such as colon, breast, stomach, lung, thyroid, esophageal, ovarian, kidney, prostate or pancreatic cancer, cesophageal, carcinoma, clear cell cancer, infiltrating ductal cancer, infiltrating lobular cancer, sequences cell carcinoma, neuroendocrine carcinoma, papillary carcinoma and Wilm's tumour.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      946 CACCGGGACCTCAAGTCCATCATCCTGATCCTGGAGGCCATCGAGAACCACAACCTC 1005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1006 GCAGACACGGTGCTCAAGATCACGGACTTCGGCCTCGCCCGCGAGTGGCACAAGACCACC 1065
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        62 TGCAATAAAACTTTGAAGATTACAGATTTTGGGTTGGCGAGGGAATGGCACAGGACCACC 121
                                                                                                                                                                                                                                                      Augustus M, Carter KC, Ebner R, Endress G, Horrigan S; Weaver \mathbb{Z}_i
                                                                                                                                                                                                                                                                                                                                     Screening for anti-neoplastic agent involves exposing cells to a chemical agent to be tested for anti-neoplastic activity, and determining a change in expression of a gene of a signature gene set
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 CACCGGGACATCAAGGCAGGAAATATTTTGCTACTTGAGAAGATAGAACATGATGACATC 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Owery Match 44.8%; Score 73.4; DB 24; Length 3454; Best Local Similarity 65.6%; Pred. No. 2.5e-11; Matches 107; Conservative 0; Mismatches 56; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human nervous system related polynucleotide SEQ ID NO 9508.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1066 AAGATGAGCGCTGCGGGGACCTACGCCTGGATGGCGCCGGAGG 1108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           122 AAAATGAGCACAGCAGCACCTATGCCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1; SEQ ID 8355; 44pp; English.
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                                          02-0CT-2000; 2000US-237295P.
02-0CT-2000; 2000US-237316P.
03-0CT-2000; 2000US-23759BP.
03-0CT-2000; 2000US-23759BP.
03-0CT-2000; 2000US-237604P.
03-0CT-2000; 2000US-23760BP.
01-WOV-2000; 2000US-23760BP.
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                                                                                                                                                                                                                                                                                                      WPI; 2002-188264/24.
                                                                                                                                                                                                                                                                      Soppet DR,
                                                                                                                                                                                                                                                        found PE,
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Best Local S
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Human; nootropic; neuroprotective; cytostatic; dermatological; virucide; immuosuppressive; antiinflammatory; anti-HTV; antibacterial; vulnerary; antiparkinsonian; antisickling; antianaemic; antiarthritic; cancer; antisickling; cerebroprotective; antiinflammatory; antialergic; antidabelic; antiolicer; anticonvisant; antiingal; antianalergic; antidabelic; antiologic; anticonvisant; antiungal; antiparasitic; cardiant; immune disorder; cardiovascular disorder; neurological disease; infection; nephrotropic; gene therapy; vaccine; ds.
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2000US-0234223.
2000US-0234274.
2000US-0234997.
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2000US-0251856
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                           14. SEP - 2000)
25 - SEP - 2000)
26 - SEP - 2000)
27 - SEP - 2000)
28 - SEP - 2000)
29 - SEP - 2000)
20 - SE
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(ABB14678-ABB18001) useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification. The nucleic acids, proteins, antibodies and (ant)agonists are useful in the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and ovarian cancer and other cancers of the adrenal gland, bone, bone marrow, breast, gastrointestinal tract, liver, lung, or urogenital; (b) immune disorders e.g. Addison's disease, allergies, autoimmune tharotic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's disease, multiple sclerosis, rheumatoid arthritis and ulcerative colitis; (c) cardiovascular disorders such as mycoardial ischaemias; (d) wound healing; (e) neurological diseases e.g. cerebral anoxia and parasitic infections.
                                                                                                                                                                                                                                                       Nucleic acids encoding 3224 human nervous system antigen polypeptides, useful for preventing, diagnosing and/or treating nervous system cancers and metastases -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4670 TTTTGCTGCTGCAGCCCATTGAGAGTGACGACATGGAGCACAAGACCCTGAAGATCACCG 4729
                                                                                                                                                                                                                                                                                                                                                                            The invention relates to novel genes (ABA11004-ABA21534) and proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    87 ATTTTGGGTTGGCGAGGGAATGGCACAGGACCACAAAATGAGCACAGCAGCACCTATG 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; immune; haematopoietic; immune/haematopoietic antigen; cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match

44.1%; Score 72.4; DB 22; Length 7084;
Best Local Similarity 70.3%; Pred. No. 7.3e-14;
Matches 97; Conservative 0; Mismatches 41; Indels 0;
                                                                                                                                                                                                                                                                                                                                    Disclosure; SEQ ID NO 9508; 1701pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 7084 BP; 1519 A; 1987 C; 2073 G; 1505 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cytostatic; gene therapy; vaccine; metastasis;
                                                                                                                                                                           Barash SC, Ruben SM;
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                    2000US-0251869.
2000US-0251989.
2000US-0251990.
2000US-0254097.
2001US-0259678.
                                                                                                                                     (HUMA-) HUMAN GENOME SCI INC
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  2000US-0251868
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2000US-0236367.
2000US-0236368.
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2000US-0236376.
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2000US-0
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Ruben SM

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amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic
activity, and can be used in gene therapy and vaccine production. (I)
proteins and polynuclectides may be used in the prevention, diagnosis and
treatment of diseases associated with inappropriate (I) expression. For
example, they may be used to treat disorders associated with decreased
expression by rectifying mutations or deletions in a patient's genome
that affect the activity of (I) by expressing inactive proteins or to
supplement the patients own production of (I). Additionally, (I)
polynuclectides may be used to produce the secreted (I), by inserting
the nucleic acids into a host cell and culturing the cell to express the
protein. (I) proteins and polynuclectides may be used to prevent,
cancers and cancer metastases of haematopoietic-derived cells. AAK64703
to AAK87694 represent human immune/haematopoietic antigen genomic
sequences from the present invention. AAK54942 to AAK54950 and AAM82169
represent sequences used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
                                     Nucleic acids encoding human immune/hematopoietic antigen polypeptides, useful for preventing, diagnosing and/or treating cancers and metastasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4670 TITIGCIGCIGCAGCCCATIGAGAGTGACGACATGGAGCCACAAGACCCTGAAGAICACG 4729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         pTK; protein tyrosine kinase; catalytic domain; c-kit; megakaryocyte; lymphocyte; amplification; primer; polymerase chain reaction; PCR; ds.
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/note= "base labelled as X in the specification"
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/note= "base labelled as X in the specification"
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/note= "base labelled as X in the specification"
                                                                                                                       Disclosure; SEQ ID NO 28728; 3071pp + Sequence Listing; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Match 44.1%; Score 72.4; DB 22; Length 7084; Local Similarity 70.3%; Pred. No. 7.3e-14; Indels 0; Mismatches 41; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 7084 BP; 1519 A; 1987 C; 2073 G; 1505 T; 0 other;
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WPI; 2001-483426/52
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The LpTKs are expressed in lymphocytic cells, as well as megakaryocytic cells. Sequencing of LpTK-4 revealed the sequences given in AAQ49751 and AAQ49755. The protein sequence corresp, to AAQ49751 is claimed (claim 7) and stated as given in the specification,
                                                                                                                                                                                                                                                                                                                                                                                                                                 pTK genes were identified using two sets of degenerative oligonucleotide primers: a first set which amplifies all pTK DNA segments (AAQA9743-44), and a second set which amplifies highly conserved sequences present in the catalytic domain of the c-kit subgroup of pTKs (AAQ49745-46). The pTK genes identified are described in AAQ49747-57 and AAR41897-02.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TTTTGCTACTTGAGAAGATAGAACATGACATCTGCAATAAAACTTTGAAGATTACAG 86
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/note= base n at position 212 is not identified
in the specification"
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/note= "base n at position 222 is not identified
in the specification"
                                                                                                                                                                                                                                                                                                            New protein tyrosine kinase genes and proteins encoded by genes are of human mega-karyocytic origin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 404;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Protein tyrosine-kinase; pTK; LpTK4; agonist; cell growth;
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69.6%; Pred. No. 2.9e-14;
tive 0; Mismatches 42; Indels
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                                                                                                                                                                                                     Scadden D;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               however is missing from the publication.
                                                                                                                                                   (NEWE-) NEW ENGLAND DEACONESS HOSPITAL.
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Conservative
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                                                                                                    92US-0826935
                                                  93WO-US00586
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                                                                                                                                                                                                                                                          WPI; 1993-320330/40.
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les 96; Conserva
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                                                    22-JAN-1993;
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05-AUG-1993
                                                                                                                                                                                                        Avraham H,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        87 ATTITGGGTTGGCGAGGGAATGGCACAGGACCACCAAAATGAGCACAGCAGGCACCTATG 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Agonist antibodies which activate specific protein tyrosine kinase(s) - also activate chimeric proteins of kinase extracellular domain and Ig constant domain, useful for studying, and therapeutic modulation of, cell growth and differentiation
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/*tag= c
/note= "base n at position 321 is not identified
in the specification"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          / Match 43.9%; Score 72; DB 16; Length 404; Local Similarity 69.6%; Pred. No. 2.9e-14; nes 96; Conservative 0; Mismatches 42; Indels
                                                                                                                                                                                                                   Lee JM, Matthews W, Tsai SP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 404 BP; 95 A; 106 C; 123 G; 77 T; 3 other;
                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 58-59; 125pp; English.
                                                                                                                                                       94US-0222616.
                                                                                                                         95WO-US04228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               223 CCTGGATGGCTCCTGAGG 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                147 CCTGGATGGCCCCAGAAG 164
                                                                                                                                                                                                                   Bennett BD, Goeddel D,
                                                                                                                                                                                     (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                WPI; 1995-366160/47.
                                                                                                                         04-APR-1995;
                                                                                                                                                       04-APR-1994;
                                                                                           12-0CT-1995.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                    Wood WI;
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